# ROYAL SOCIETY OPEN SCIENCE

#### rsos.royalsocietypublishing.org

# Research



**Cite this article:** Lim AN, De La Guerra JA, Blumstein DT. 2016 Sea anemones modify their hiding time based on their commensal damselfish. *R. Soc. open sci.* **3**: 160169. http://dx.doi.org/10.1098/rsos.160169

Received: 8 March 2016 Accepted: 27 July 2016

Subject Category:

Biology (whole organism)

Subject Areas: behaviour

#### **Keywords:**

hiding time, antipredator, defence, dynamic mutualism, anemones, anemonefish

#### Author for correspondence:

Daniel T. Blumstein e-mail: marmots@ucla.edu

Electronic supplementary material is available at http://dx.doi.org/10.1098/rsos.160169 or via http://rsos.royalsocietypublishing.org.



# Sea anemones modify their hiding time based on their commensal damselfish

# Alexandra N. Lim, Justin A. De La Guerra and Daniel T. Blumstein

Department of Ecology and Evolutionary Biology, University of California, 621 Young Drive South, Los Angeles, CA 90095-1606, USA

🔟 DTB, 0000-0001-5793-9244

Animals often retreat to refugia when alarmed and the time they spend hiding reflects an economic decision that trades off reducing predation risk with other beneficial activities. Typically, refugia such as burrows are static, but some refugia are dynamic. For species with defensive mutualisms, hiding might be contingent on their mutualist's behaviour. We disturbed and quantified hiding time in magnificent sea anemones, *Heteractis magnifica*, and their associated domino damselfish, *Dascyllus trimaculatus*. We found that sea anemone hiding behaviour was dependent on the number and behaviour of their commensal fish: anemones emerged sooner when they had more associated fish and faster returning fish. Together, these results demonstrate that hiding behaviour can be influenced by the behaviour of a commensal; such dynamic mutualisms may be found in other systems.

## 1. Introduction

Prey that do not escape predators or that suffer non-fatal injuries incur a cost, and there are a variety of complex antipredator adaptations to reduce these costs [1]. For example, many animals hide in protective refuges for variable lengths of time. Fiddler crabs, *Uctea lactea perplexa*, retreat to burrows, and polychaete worms, *Serpula vermicularis*, retreat into their calcareous tubes [2]. Variation in hiding, like many other antipredator behaviours, has an economic logic. The optimal hiding time model [3] suggests that antipredator hiding time decisions are based on a balance between predation risk and the cost of refuge use. Once hidden, prey can emerge when the predator is still present and be killed, or stay in the refuge long after the predator is absent and lose foraging opportunities [4].

A defining characteristic of previous studies has been that they involved static hiding systems where organisms sought shelter in unchanging refugia. Examples of static terrestrial

© 2016 The Authors. Published by the Royal Society under the terms of the Creative Commons Attribution License http://creativecommons.org/licenses/by/4.0/, which permits unrestricted use, provided the original author and source are credited.

refugia include trees, cliff faces, burrows, thick vegetation and rock talus [5]. However, there are some systems where shelters are dynamic rather than static. One such system is the anemone–anemonefish mutualism. In these systems, host anemones create dynamically changing shelters for anemonefish, whereas fish defend their host anemones from predation [6].

Sea anemones avoid predators by tensing or withdrawing their tentacles into the upper column of their body cavity and effectively hiding [7]. When closed, commensal anemonefish hide among rocks or below the skirt of their host anemone. Unlike other static systems, the antipredator hiding behaviour of anemones and anemonefish are highly dependent on one another and can be described as a dynamic protection mutualism. Their dynamic behaviour makes this an excellent system to ask whether variation in fish behaviour affects anemone hiding.

## 2. Material and methods

#### 2.1. Study site and subjects

We studied magnificent sea anemones, *Heteractis magnifica*, attached to individually mapped coral bommies, and domino damselfish, Dascyllus trimaculatus, between 22 January and 30 January 2016 in Opunohu and Ha'apita Beaches, Mo'orea, French Polynesia (17.53°S, 149.83°W). All Opunohu anemones (n = 24) had commensal anemonefish, whereas Ha'apita anemones (n = 12) had none. When bommies had more than one anemone, we randomly selected one of the presumed clones for study [8]. We counted the total number of anemonefish associated with a bommie and categorized them into three size classes: small (less than 2 cm), medium (2-5 cm) and large (more than 5 cm) maximum length. Anemones fire either or both offensive or defensive nematocysts in response to biological stimulation [9]. We therefore collected an abundant macroalgae, Turbinaria ornata, and systematically disturbed the anemones and their associated fish by swiping the anemone with algal thalli seven times. The first researcher (A.N.L.) disturbed the anemone and retreated 1.5 m away; a distance that pilot observations showed would not interfere with fish behaviour. Anemones responded to rubbing by tensing and retracting their tentacles and occasionally lifting their skirt off the substrate. The second researcher, at a distance of 1.5 m, video recorded (Fujifilm FinePix XP80 or Olympus Tough digital cameras) all experiments 30 s prior to disturbance until the anemone was fully relaxed and all anemonefish had returned. We measured, from the video, the latency to initial anemone movement as the time from disturbance until first movement of tentacles, and the latency to total anemone relaxation as the time from the disturbance until the subject was freely moving all tentacles and its skirt was completely relaxed. We also measured latency to first fish return and latency to all fish return as the time from when all fish hid until the first fish and then all fish returned to the anemone. All anemones were 1-2 m deep. To obtain a better estimate of the repeatability of hiding time and fish counts, we resampled 23 marked Opunohu individuals three times—once every other day.

#### 2.2. Statistical analyses

Using the full dataset of the day one experimental manipulations on each anemone, we fitted general linear models (GLMs) to explain variation in the latency to initial anemone movement following disturbance as well as the latency to full anemone relaxation as a function of total number of fish. Using the Opunohu dataset of day one experimental manipulations, we fitted linear models to explain anemone behaviour as a function of total number of fish, latency to first anemonefish return and latency to all anemonefish return.

To quantify the consistency of our estimates of both fish counts and of hiding time, we fitted a series of linear mixed-effects models on the Opunohu data. We first fitted a random intercept model where we explained variation in the number of fish or hiding time as a function only of anemone ID. This permitted us to calculate the intraclass correlation coefficient, a measure of repeatability for the number of fish, latency to anemone initial movement and latency to total relaxation. We then fitted a random intercept model and a random slope model with a set of fixed effects. In all cases, we included trial number, total number of fish, latency to first fish return and latency to all fish return as fixed effects. We compared the mixed models to a GLM with no random effects using likelihood ratio tests (LRTs).

We  $log_{10}$ -transformed total number of fish, latency to initial anemone movement, latency to total anemone relaxation, latency to first fish return and latency to all fish return. Analyses were conducted in R v. 3.2.3 [10]. We used Deducer [11] to fit the GLMs, the R packages lme4 [12] for the mixed-effects models and lmeTest [13] to calculate *p*-values for the mixed models. For the GLMs, we report type III

sums of squares and adjusted  $R^2$ -values. We calculated likelihood ratio tests using lme4 and RSLsim [14]. Our residuals were approximately normally distributed. We plotted residuals versus fitted values and found two to three outliers. While there was no reason to remove them, we did so, and recalculated the models. The overall model fit improved after removing outliers, but there was no change in the general conclusions from the models with the outliers. We thus report the results of the dataset including statistical outliers.

### 3. Results

#### 3.1. Latency to first movement following disturbance

Anemones varied in their number of commensal fish. While anemones at Opunohu had at least some commensal fish, anemones at Ha'apita had none. Using the full dataset, we found that anemones with more commensal anemonefish first moved sooner after experimental disturbance (estimate = -0.202, p = 0.0187; figure 1*a*). This model explained 12.7% of variation in anemone hiding time.

Focusing only on the subset of anemones with commensal fish at Opunohu, and after explaining variation accounted for by total number of fish (estimate = -0.206, p = 0.319), latency to first fish return (estimate = -0.567, p = 0.0260) and latency to all fish return (estimate = 0.337, p = 0.0283), we found that anemones with a shorter latency to first fish return had a longer latency to initial tentacle movement (figure 1*b*). Anemones with fish that all returned sooner, first moved their tentacles sooner (figure 1*c*). This model explained 15.4% of variation in anemone hiding time (p = 0.107).

For Oponohu data, we explored how fish size influenced hiding time by modelling latency to first move as a function of either the number of small (estimate = -0.158, p = 0.158), medium (estimate = -0.225, p = 0.0706) or large (estimate = -0.578, p = 0.002) fish. Thus, the number of large fish best explained variation in anemone hiding behaviour.

#### 3.2. Latency to full relaxation following disturbance

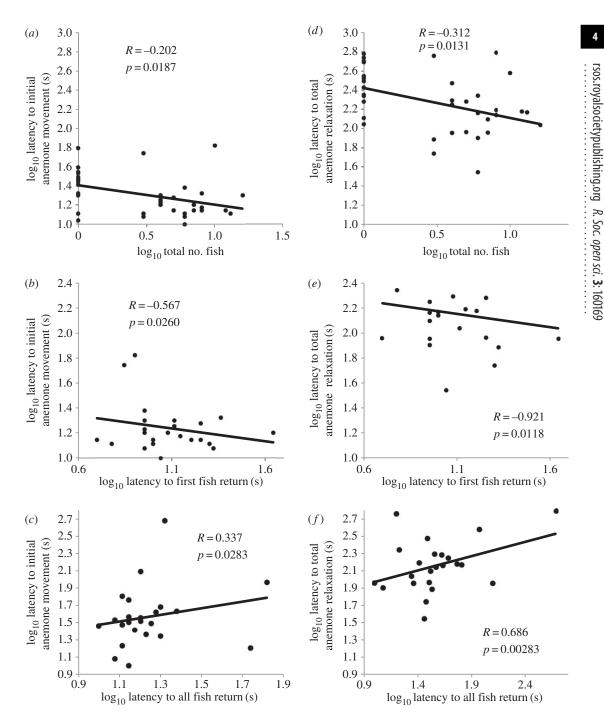
In the full dataset, we found that anemones with more commensal anemonefish fully relaxed sooner (estimate = -0.312, p = 0.0131; figure 1*d*). The model explained 14.33% of variation in anemone hiding time.

Focusing only on the subset of anemones with commensal fish at Opunohu, and after explaining variation accounted for by total number of fish (estimate = -0.257, p = 0.375), latency to first fish return (estimate = -0.921, p = 0.0118), and latency to all fish return (estimate = 0.686, p = 0.00283), we found that anemones with shorter latency to first fish return had longer latency to relax (figure 1*e*). Anemones with shorter latency to all fish return fully relaxed sooner (figure 1*f*). This model explained 30.4% of variation in anemone hiding time (p = 0.0194).

For Oponohu data, we explored how fish size influenced hiding time by modelling latency to total relaxation as a function of either the number of small (estimate = -0.276, p = 0.0918), medium (estimate = -0.321, p = 0.0805) or large (estimate = -0.748, p = 0.0066) fish. The number of large fish best explained variation in anemone latency to total relaxation.

#### 3.3. Consistency in anemone response behaviour

Fish counts were highly repeatable over 3 days (ICC = 0.84). After accounting for variation explained by our fixed factors, individual anemones had significantly different latencies to first movement (p = 0.0291; figure 2a and electronic supplementary material, table S1a) but they did not differ in their slopes (p = 0.0556; electronic supplementary material, table S1b). This random intercept model on latency to first movement was superior to a GLM with no random effects (LRT = 17.6, p = 2.2 × 10<sup>-16</sup>; electronic supplementary material, table S2a). Similarly, after accounting for variation explained by our fixed factors, individual anemones had significantly different latencies to total relaxation (p = 0.000488; figure 2b and electronic supplementary material, table S1a) but they did not differ in their slopes (p = 6.97 × 10<sup>-15</sup>; electronic supplementary material, table S1b). This random intercept model on latency to total relaxation was superior to a GLM with no random effects (LRT = 7.10, p = 0.0023; electronic supplementary material, table S2b).

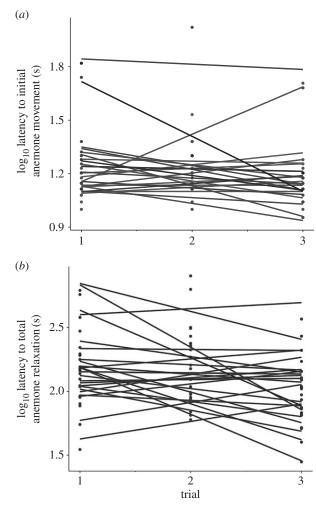


**Figure 1.** Log<sub>10</sub> latency for initial movement following disturbance as a function of (*a*)  $\log_{10}$  total number of fish, (*b*)  $\log_{10}$  time for first anemonefish to return and (*c*)  $\log_{10}$  time for all fish return. Log<sub>10</sub> latency to total anemone relaxation following disturbance as a function of (*d*)  $\log_{10}$  total number of fish, (*e*)  $\log_{10}$  time for first anemonefish to return and (*f*)  $\log_{10}$  time for all fish return.

### 4. Discussion

We found that variation in anemone behaviour was contingent on number and behaviour of anemonefish on two timescales: both the short term and the long term. Anemones moved and fully relaxed sooner when associated with fish with shorter latencies to return. Likewise, anemones associated with more commensal fish moved and reached full relaxation sooner following disturbance. Thus, variation in anemonefish behaviour explains variation in anemone hiding time.

Strictly, because all anemones at the Opunohu site were associated with fish, while none at the Ha'apiti site were, it was not possible to statistically isolate any effects of site on anemone response.



**Figure 2.** Repeatability of (*a*) anemone latency to first movement and (*b*) latency to total relaxation over three separate experimental disturbances. A random intercept model best explained variation in hiding time.

We do not know why there were no fish associated with the anemones at the Ha'apiti site. Both the anemones, and the site itself appeared healthy. Future studies, at more sites, will be required to better understand the effects of site variation on anemone antipredator behaviour.

Anemones vary in their size, and this could influence the latency it takes them to relax. A set of complementary analyses (electronic supplementary material, table S3) revealed that for almost all of the analyses, the addition of maximum anemone diameter (a measure of size) as a covariate either failed to improve the model, had no significant effect on latency to relax, or eliminated the significance of the models entirely. From this, we conclude that anemone size has a limited effect on the time it takes them to relax following disturbance.

These results demonstrate for the first time that variation in hiding behaviour can be influenced by the variable behaviour of a commensal species. While there are a number of well-studied protective mutualisms including ants and their plants, and squid and their bioluminescent bacteria, no prior studies have analysed the dynamics of hiding in the protected species [15]. Interestingly, it seems that the number of large fish may drive this relationship, because our analysis suggests that only the number of large fish explained variation in hiding time. A previous study found that anemone growth rate depended on the size and number of anemonefish and that anemones with more large fish had higher growth rates [16]. This dynamic protective mutualism appears more complex, because the benefits and costs may vary as a function of fish size to both the anemone and the fish themselves.

Dynamic mutualisms may exist in other systems. Future studies should evaluate various marine and terrestrial mutualisms to better understand how defensive behaviour can dynamically change depending on the behaviour of a mutualistic partner.

5

Ethics. All experiments were conducted under permits issued by the Government of French Polynesia (protocole d'accueil issued on 23 October 2015), and under UCLA IACUC protocol no. 2000-147).

Data accessibility. The dataset supporting this article is provided as electronic supplementary material.

Authors' contributions. A.N.L., J.A.D.L.G. and D.T.B. conceived the idea. A.N.L. and J.A.D.L.G. collected the field data and A.N.L., J.A.D.L.G. and D.T.B. analysed the data and wrote and edited the manuscript.

Competing interests. We declare we have no competing interests.

Funding. We thank the UCLA Office of Instructional Development, and the Department of Ecology and Evolutionary Biology for partial support.

Acknowledgements. We thank the UC Berkeley Gump South Pacific Research Station for logistical support, Keith Seydel, Caitlin Fong, Tiffany Armenta and Peggy Fong for assistance, and two anonymous reviewers for comments on a previous version of the manuscript.

## References

- Cooper WE, Blumstein DT. 2015 Escape behavior: importance, scope, and variables. In *Escaping from* predators: an integrative view of escape decisions (eds WE Cooper, DT Blumstein), pp. 3–11. Cambridge, UK: Cambridge University Press.
- Dill LM, Fraser AHG. 1996 The worm re-turns: hiding behavior of a tube-dwelling marine polychaete. Serpula vermicularis. Behav. Ecol. 8, 186–193. (doi:10.1093/beheco/8.2.186)
- Martín L, Lopez P. 2015 Hiding time in refuge. In *Escaping from predators: an integrative view of escape decisions* (eds WE Cooper, DT Blumstein), pp. 227–253. Cambridge, UK: Cambridge University Press.
- Sih A. 1992 Prey uncertainty and the balancing of antipredator and feeding needs. Am. Nat. 139, 1052–1069. (doi:10.1086/285372)
- Cowlishaw G. 1997 Refuge use and predation risk in a desert baboon population. *Anim. Behav.* 54, 241–253. (doi:10.1006/anbe.1996.0466)

- Holbrook SJ, Schmitt RJ. 2005 Growth, reproduction and survival of a tropical sea anemone (*Actiniaria*): benefits of hosting anemonefish. *Coral Reefs* 24, 67–73. (doi:10.1007/s00338-004-0432-8)
- Haag E, Dyson K. 2014 Trade-off between safety and feeding in the sea anemone *Anthopleura aureoradiata*. *New Zeal. J. Mar. Fresh.* 48, 540–546. (doi:10.1080/00288330.2014.915858)
- Francis L. 1973 Clone specific segregation in the sea anemone *Anthropleura elegantissima*. *Biol. Bull.* 144, 64–72. (doi:10.2307/1540147)
- Fautin DG. 2009 Structural diversity, systematics, and evolution of cnidae. *Toxiconomy* 54, 1054–1064. (doi:10.1016/j.toxicon.2009.02.024)
- R Core Team. 2015 R: a language and environment for statistical computing. Vienna, Austria: R Foundation for Statistical Computing. See https://www.Rproject.org/.
- Fellows I. 2012 Deducer: a data analysis GUI for R. J. Stat. Softw. 49, 1–15. (doi:10.18637/jss.v049.i08)

- Bates D, Maechler M, Bolker B, Walker S. 2015 Fitting linear mixed-effects models using lme4. J. Stat. Softw. 67, 1–48. (doi:10.18637/jss.v067.i01)
- Kuznetsova A, Brockhoff PB, Christensen RHB. 2015 ImerTest: tests in linear mixed effects models.
  R package version 2.0-29. See https://CRAN.Rproject.org/package=ImerTest.
- Scheipl F, Greven S, Kuechenhoff H. 2008 Size and power of tests for a zero random effect variance or polynomial regression in additive and linear mixed models. *Comput. Stat. Data Anal.* 52, 3283–3299. (doi:10.1016/j.csda.2007.10.022)
- Baskett CA, Schemske DW. 2015 Evolution and genetics of mutualism. In *Mutualism* (ed. JL Bronstein), pp. 77–93. Oxford, UK: Oxford University Press.
- Porat D, Chadwick-Furman NE. 2004 Effects of anemonefish on giant sea anemones: expansion behavior, growth, and survival. *Hydrobiol.* 530, 513–520. (doi:10.1007/s10750-004-2688-y)